

FIG. 1

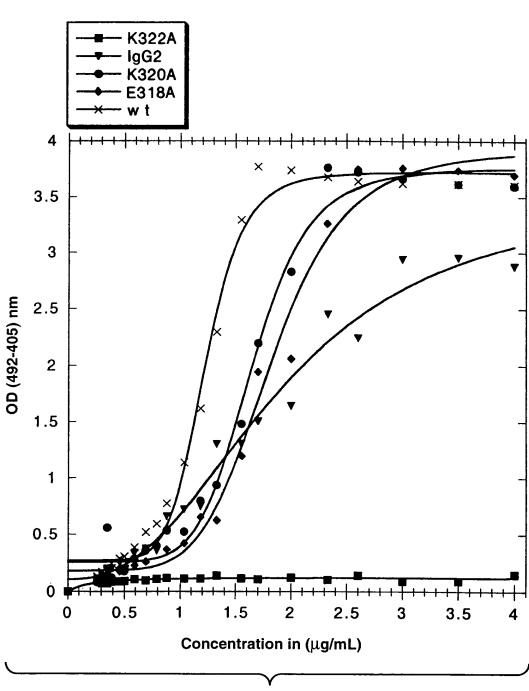


FIG. 2

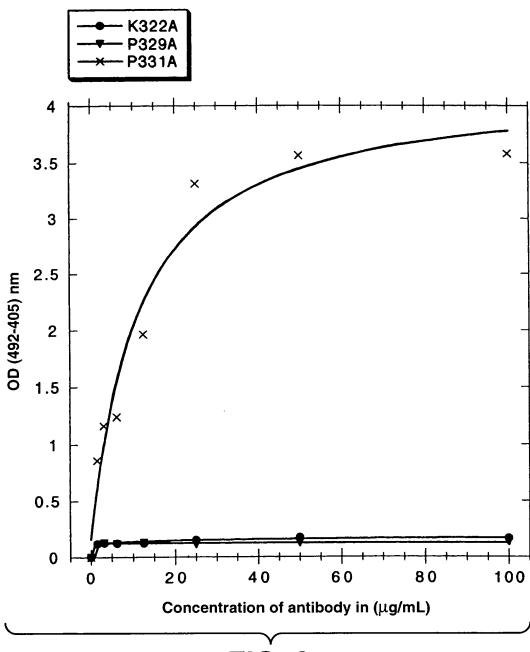


FIG. 3

4/26

GVPSRFSGSG SGTASVVCLL THOGLSSPVT LIYAASYLES EKHKVYACEV IFPPSDEQLK QQKPGKAPKL KRTVAAPSVF STLTLSKADY ITCRASKPVD GEGDSYLNWY SLQPEDFATY YCQQSHEDPY TFGQGTKVEI GNSQESVTEQ DSKDSTYSLS LSASVGDRVT **OWKVDNALQS** DIQLTQSPSS SGTDFTLTIS NNFYPREAKV KSFNRGEC

(E27) - Light Chain

F/G. 4A

(E27) - Heavy Chain

SIKYSGETKY NPSVKGRITI PLAPSSKSTS TYICHVNHKP SHEDPEVKFN SKAKGQPREP YSKLTVDKSR SASTKGPSVF TVPSSSLGTQ IAVEWESNGQ PENNYKTTPP VLDSDGSFFL PEVTCVVDV ALPAPIEKTI SGYSWNWIRQ APGKGLEWVA KEYKCKVSNK PKDTLMISRT WGQGTLVTVS SGLYSLSSVV TVLHQDWLNG HYFGHWHFAV VHTFPAVLQS GPSVFLFPPK SCAVSGYSIT TAVYYCARGS SWNSGALTSG PPCPAPELLG NSTYRVVSVL CLVKGFYPSD TOKSLSLSPG EVQLVESGGG LVQPGGSLRL LOMNSLRAED KDYFPEPVTV PKSCDKTHTC AKTKPREEOY EMTKNOVSLT VMHEALHNHY WOOGNVFSCS SRDDSKNTFY GGTAALGCLV SNTKVDKKVE WYVDGVEVHN QVYTLPPSRE

FIG. 4B

5 / 26

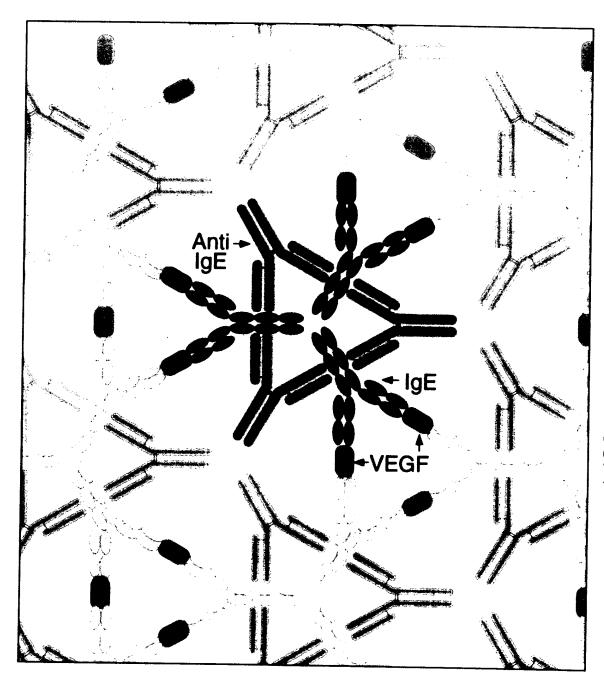
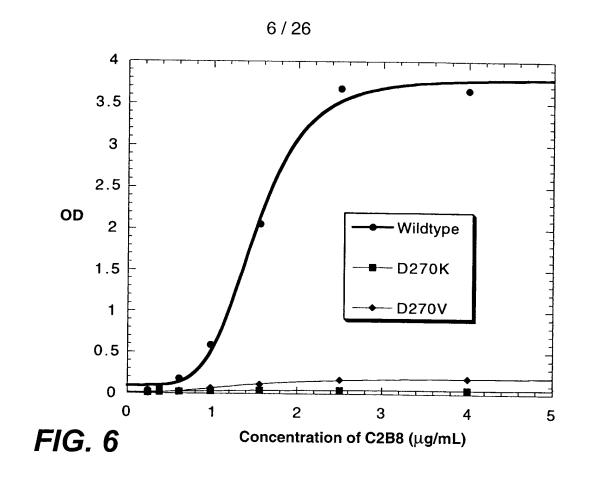


FIG. 5



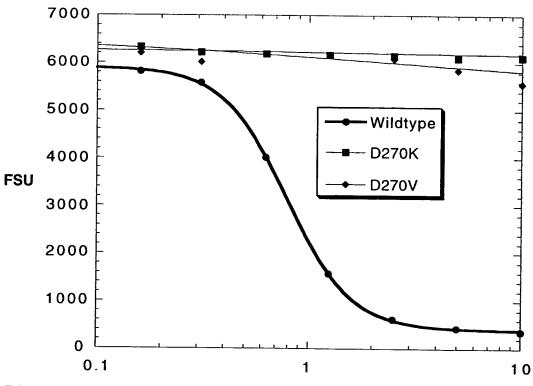


FIG. 7 Log of Concentration of C2B8 (μg/mL)

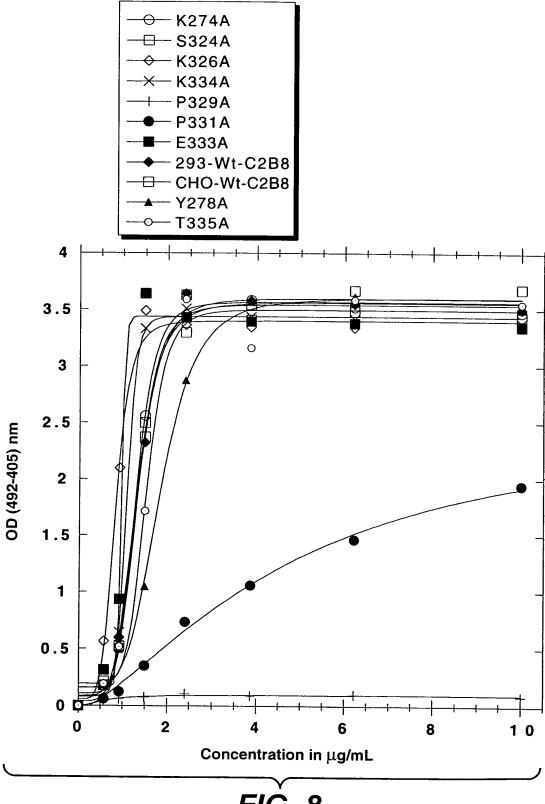


FIG. 8

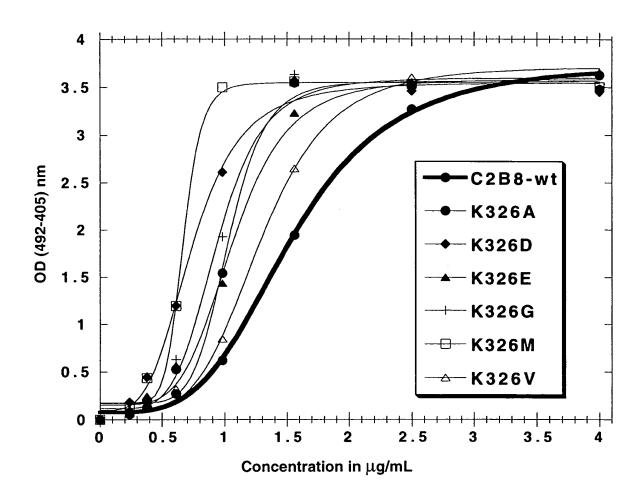


FIG. 9

P1726R1P1

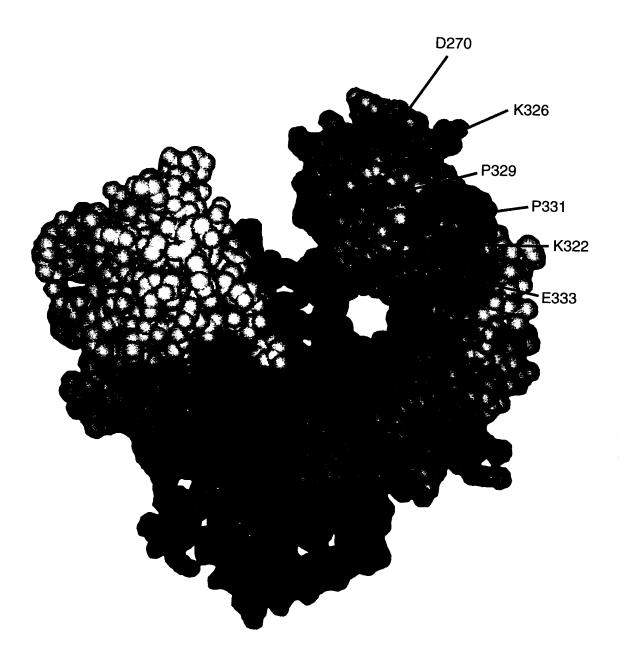


FIG. 10

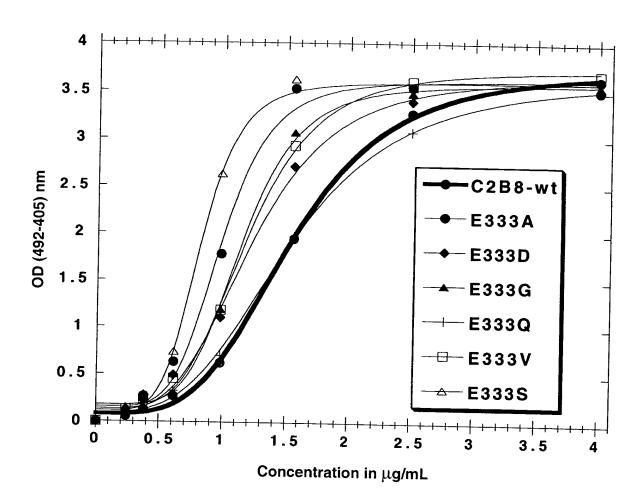
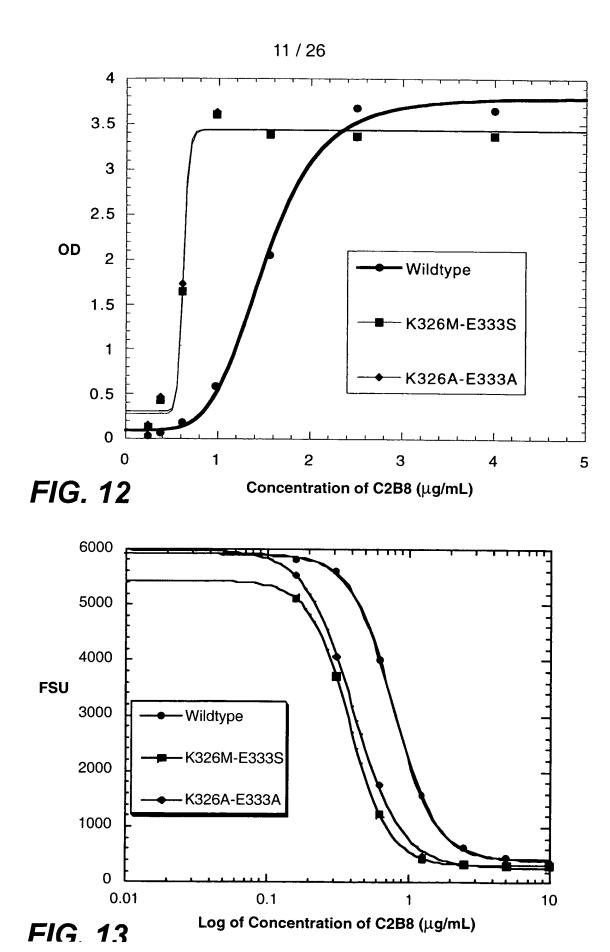


FIG. 11



12/26

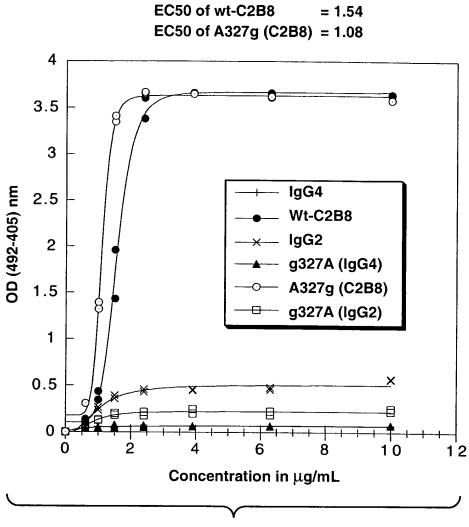
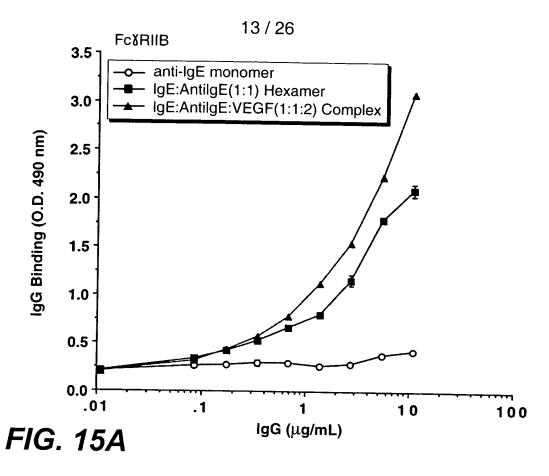
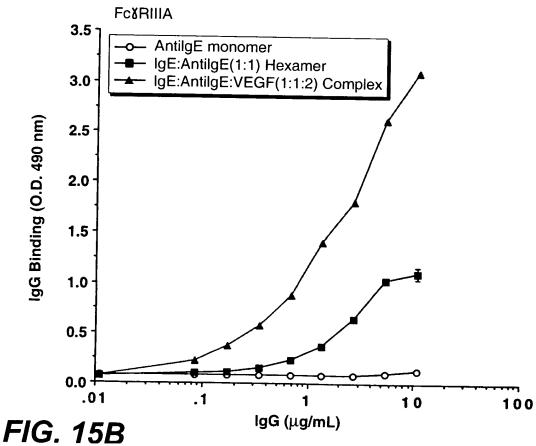
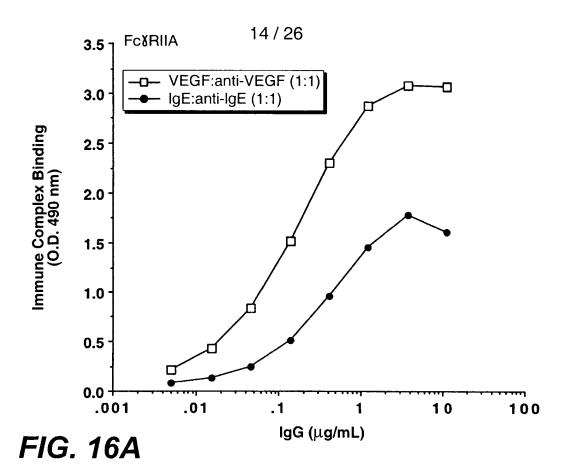
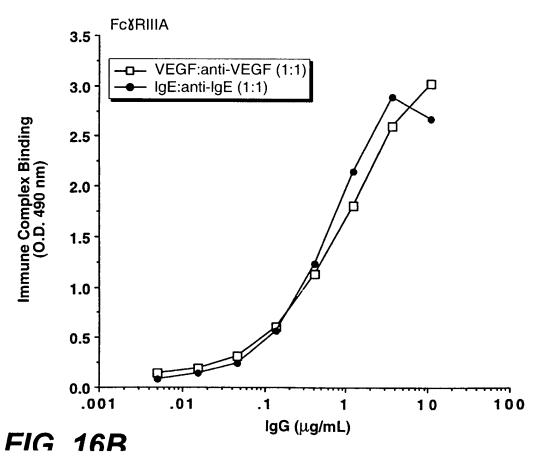


FIG. 14

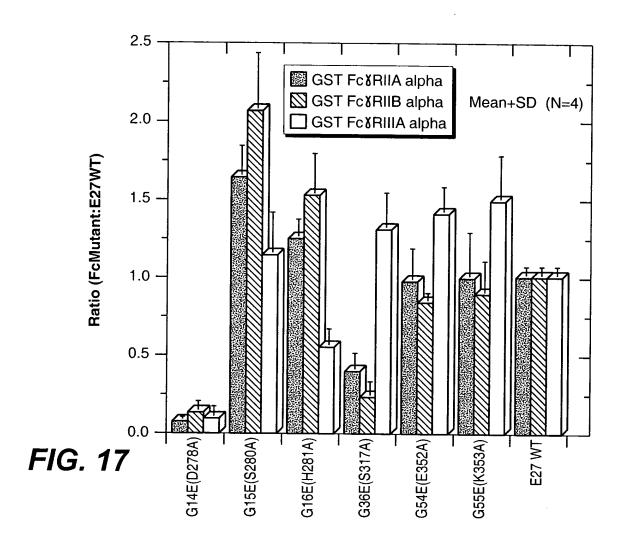


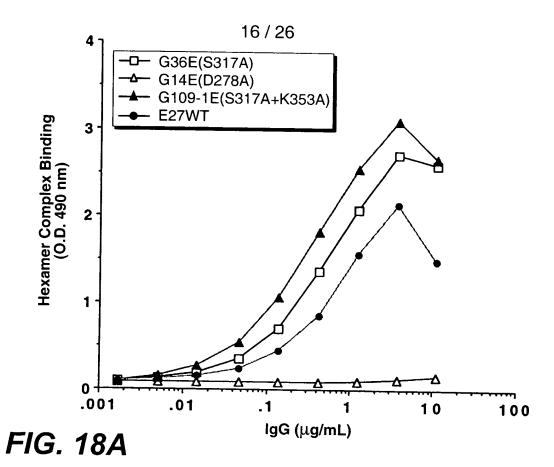






15 / 26





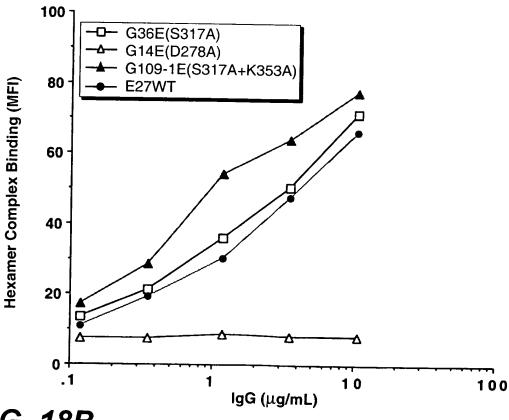
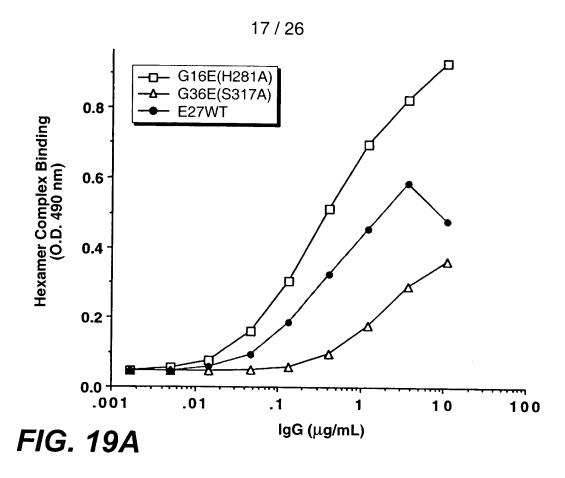
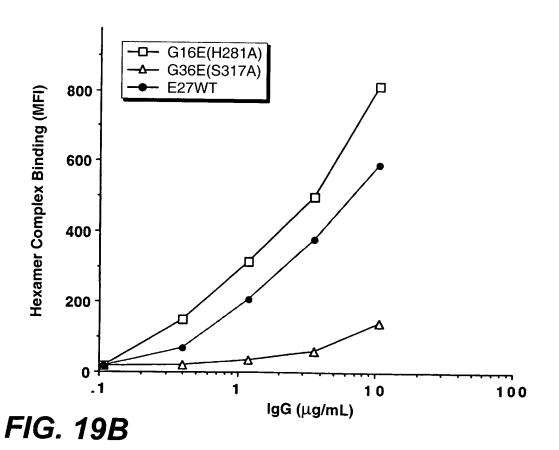


FIG. 18B





18 / 26

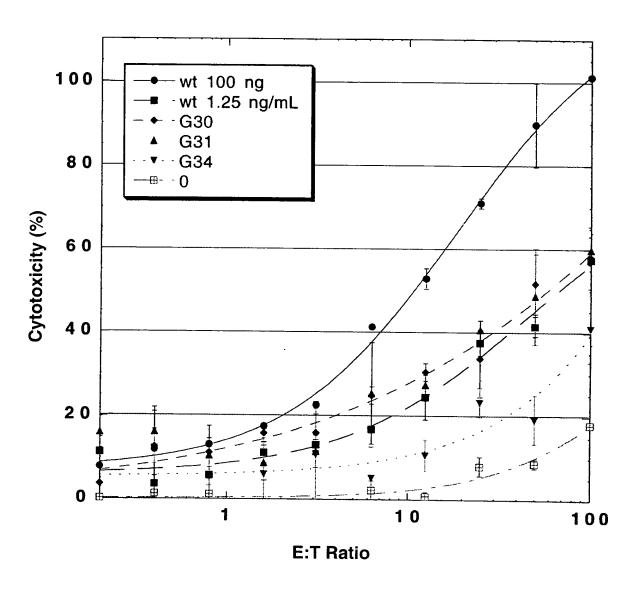


FIG. 20

19/26

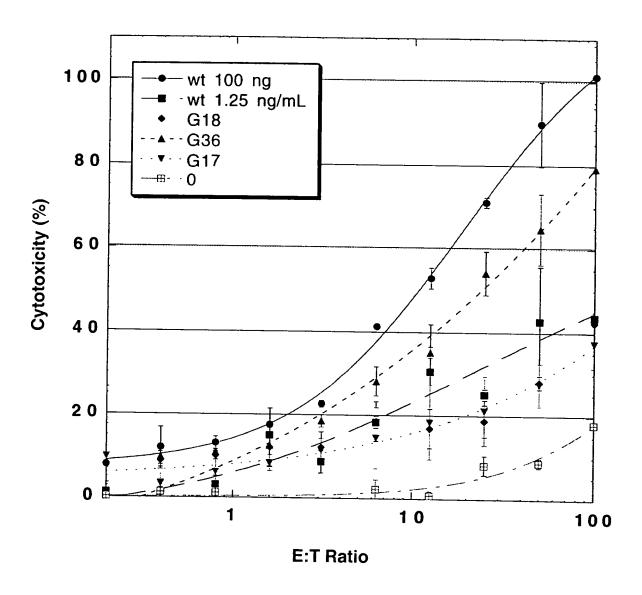


FIG. 21

P1726R1P1

humIgG1 humIgG2 humIgG3 humIgG4 murIgG1 murIgG2A murIgG2B murIgG3	230 240 250 260 270 PAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYV PAP-PVAGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVQFNWYV PAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVQFKWYV PAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVTVPEVSSVFIFPPKPKDVLTITLTPKVTCVVVDISKDDPEVQFSWFV PAPNLLGGPSVFIFPPKIKDVLMISLSPIVTCVVVDVSEDDPDVQISWFV PAPNLEGGPSVFIFPPNIKDVLMISLTPKVTCVVVDVSEDDPDVQISWFV PPGNILGGPSVFIFPPKPKDALMISLTPKVTCVVVDVSEDDPDVHVSWFV
humIgG1 humIgG2 humIgG3 humIgG4 murIgG1 murIgG2A murIgG2B murIgG3	280 290 300 310 320 DGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALP DGVEVHNAKTKPREEQFNSTFRVVSVLTVVHQDWLNGKEYKCKVSNKGLP DGVEVHNAKTKPREEQFNSTFRVVSVLTVLHQDWLNGKEYKCKVSNKALP DGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKGLP DDVEVHTAQTQPREEQFNSTFRSVSELPIMHQDCLNGKEFKCRVNSAAFP NNVEVHTAQTQTHREDYNSTLRVVSALPIQHQDWMSGKEFKCKVNNKDLP NNVEVHTAQTQTHREDYNSTIRVVSHLPIQHQDWMSGKEFKCKVNNKDLP DNKEVHTAWTQPREAQYNSTFRVVSALPIQHQDWMRGKEFKCKVNNKALP
humIgG1 humIgG2 humIgG3 humIgG4 murIgG1 murIgG2A murIgG2B murIgG3	330 340 350 360 370 APIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAV D L APIEKTISKTKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAV APIEKTISKTKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAV SSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAV APIEKTISKTKGRPKAPQVYTLPPPKEQMAKDKVSLTCMITDFFPEDITV APIERTISKPKGSVRAPQVYVLPPPEEEMTKKQVTLTCMVTDFMPEDIYV SPIERTISKPKGLVRAPQVYTLPPPAEQLSRKDVSLTCLVVGFNPGDISV APIERTISKPKGRAQTPQVYTIPPPREQMSKKKVSLTCLVTNFFSEAISV
humIgG1 humIgG2 humIgG3 humIgG4 murIgG1 murIgG2A murIgG2B murIgG3	380 390 400 410 420 EWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMH EWESNGQPENNYKTTPPMLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMH EWESSGQPENNYNTTPPMLDSDGSFFLYSKLTVDKSRWQQGNIFSCSVMH EWZSNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMH EWQWNGQPAENYKNTQPIMDTDGSYFVYSKLNVQKSNWEAGNTFTCSVLH EWTNNGKTELNYKNTEPVLDSDGSYFMYSKLRVEKKNWVERNSYSCSVVH EWTSNGHTEENYKDTAPVLDSDGSYFIYSKLNMKTSKWEKTDSFSCNVRH EWERNGELEQDYKNTPPILDSDGTYFLYSKLTVDTDSWLQGEIFTCSVVH
humIgG1 humIgG2 humIgG3 humIgG4 murIgG1 murIgG2A murIgG2B murIgG3	430 440 EALHNHYTQKSLSLSPGK EALHNHYTQKSLSLSPGK EALHNRFTQKSLSLSPGK EALHNHYTQKSLSLSLGK EGLHNHHTEKSLSHSPGK EGLHNHHTTKSFSRTPGK EGLKNYYLKKTISRSPGK EALHNHHTQKNLSRSPGK

P1726R1P1

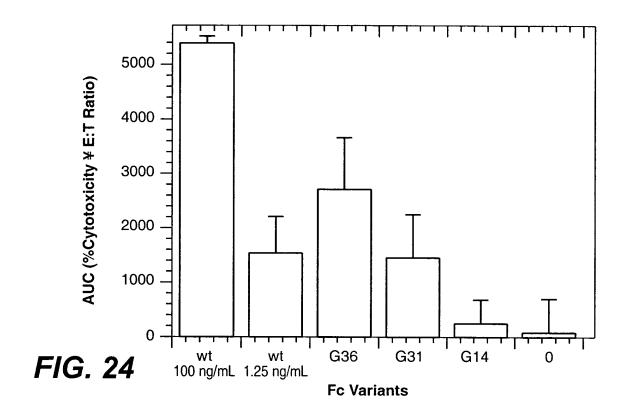
21 / 26

Percent Identity Among Fc Sequences

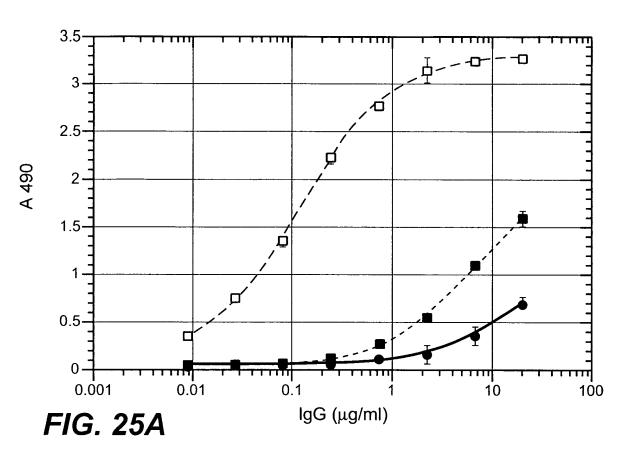
		1	2	3	4	5	6	7	8
1.	humIgG1	-	94	94	94	64	66	63	68
2.	humIgG2		-	93	92	65	63	60	67
3.	humIgG3			-	91	64	64	61	67
4.	humIgG4				_	62	64	61	64
5.	murIgG1					_	65	61	67
6.	murIgG2A						-	77	70
7.	murIgG2B							_	68
8.	murIqG3								_

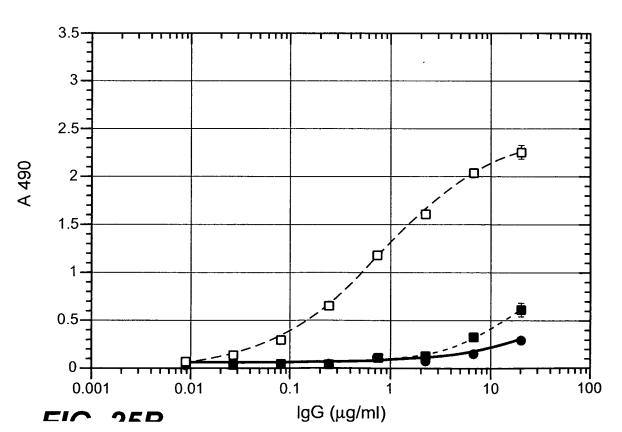
FIG. 22B

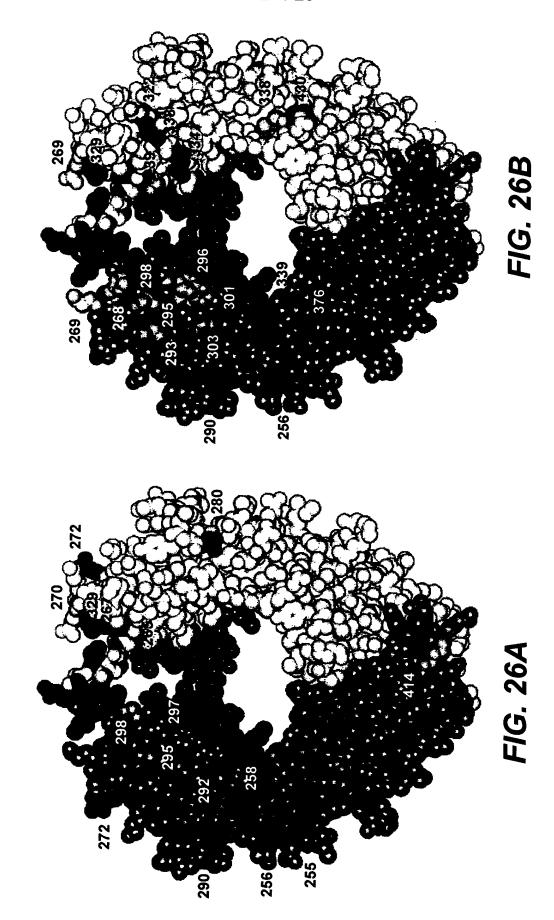
humIgG1 humIgG2 humIgG3 humIgG4	PAP-PVA PAPELLG	GPSVFLFPPK GPSVFLFPPK	PKDTLMISRT PKDTLMISRT	PEVTCVVVDV:	270 SHEDPEVKFNWYV SHEDPEVQFNWYV SHEDPEVQFKWYV SQEDPEVQFNWYV * * *
humIgG1 humIgG2 humIgG3 humIgG4	DGVEVHNI DGVEVHNI	AKTKPREEQF AKTKPREEQF	NSTFRVVSVL NSTFRVVSVL	.TVVHQDWLNG .TVLHQDWLNG	320 KEYKCKVSNKALP KEYKCKVSNKGLP KEYKCKVSNKALP KEYKCKVSNKGLP
humIgG1	330 APIEKTIS	340 SKAKGQPREP		360 EMTKNQVSLTO	370 CLVKGFYPSDIAV
humIgG2 humIgG3 humIgG4	APIEKTIS	KTKGQPREP(QVYTLPPSRE QVYTLPPSRE	EMTKNQVSLTO	CLVKGFYPSDIAV CLVKGFYPSDIAV CLVKGFYPSDIAV
humIgG1 humIgG2 humIgG3 humIgG4	EWESNGQI EWESSGQI	PENNYKTTPPI PENNYNTTPPI PENNYKTTPPI	MLDSDGSFFL MLDSDGSFFL	YSKLTVDKSRV YSKLTVDKSRV	420 VQQGNVFSCSVMH VQQGNVFSCSVMH VQQGNIFSCSVMH VQQGNVFSCSVMH VQEGNVFSCSVMH
humIgG1 humIgG2 humIgG3 humIgG4	EALHNHYT EALHNRFT	440 COKSLSLSPGI COKSLSLSPGI COKSLSLSPGI COKSLSLSLGI	ë Fl	G. 23	



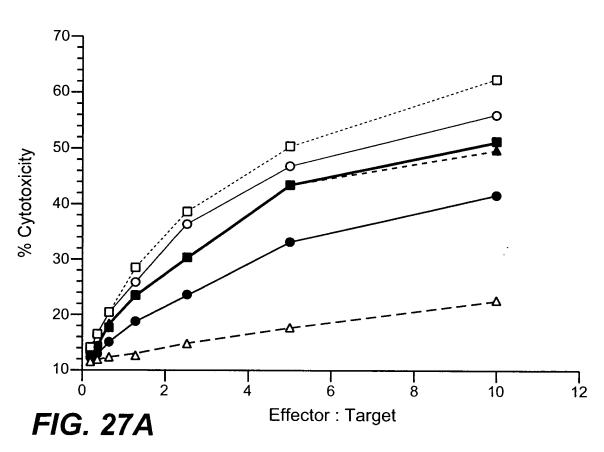
23 / 26

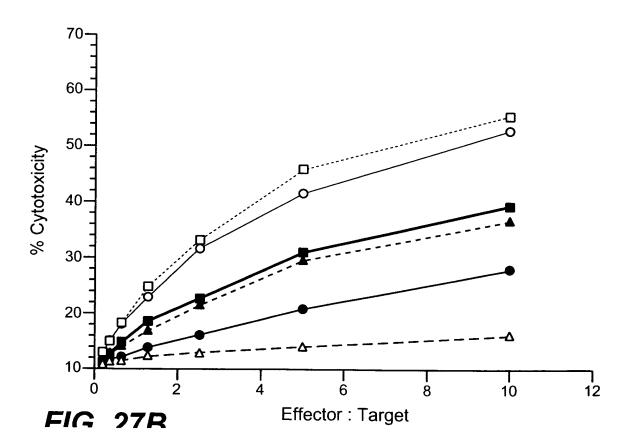












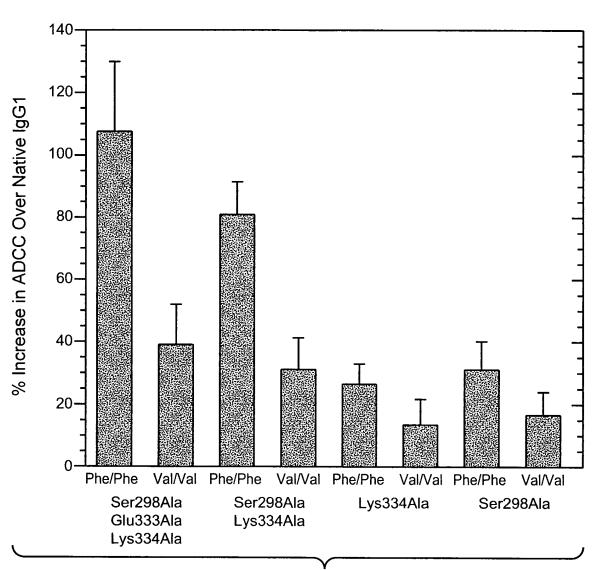


FIG. 27C